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Applicant(s): Edwin Madison et al.

Nucleic Acid Molecules Encoding A Transmembrane Serine Protease 7, The Encoded Polypeptides And Methods Based

Thereon

Domain organization and amino acid sequence of MTSP7

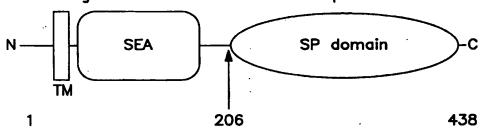


FIG. 1A

MAKEROTER	10	20				60				
MMYTPVEFSEAEFSRAEYQRKQQFWDSVRLALFTLAIVAIIGIAIGIVTHFVVEDDKSFY										
	70	80	90	100	110	120				
YLASFKVTNIKYKENYGIRSSREFIERSHQIERMMSRIFRHSSVGGRFIKSHVIKLSPDE										
;	130	140	150	160	170	180				
QGVDILIVLIFRYPSTDSAEQIKKKIEKALYQSLKTKQLSLTINKPSFRLTPIDSKKMRN										
:	190	200 1	210	220	230	240				
190 200 1 210 220 230 240 LLNSRCGIRMTSSNMPLPASSSTQRIVQGRETAMEGEWPWQASLQLIGSGHQCGASLISN										
;	250	260	270	280	290	300				
TWLLTAAHCFWKNKDPTQWIATFGATITPPAVKRNVRKIILHENYHRETNENDIALVQLS										
,	310	320	330	340	350	360				
TGVEFSNIVQRVCLPDSSIKLPPKTSVFVTGFGSIVDDGPIQNTLRQARVETISTDVCNR										
,	370	380	390	400	410	420				
KDVYDGLITPGMLCAGFMEGKIDACKGDSGGPLVYDNHDIWYIVGIVSWGQSCALPKKPG										
	430									
VYTRVTKYRDWIASKTGM*										

↓ = protease cleavage site

FIG. 1B

Replacement Sheet

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FIG. 1C

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970	980	990	1000	1010	1020						
CAAATATAGTCCAGAGAGTTTGCCTCCCAGACTCATCTATAAAGTTGCCACCTAAAACAA GTTTATATCAGGTCTCTCAAACGGAGGGTCTGAGTAGATATTTCAACGGTGGATTTTGTT											
GTTTATATC	GGTCTCTCA	VACGGAGGG	CTGAGTAGA:	PATTTCAACGO	TGGATTTTGTT						

- GTGTGTTCGTCACAGGATTTGGATCCATTGTAGATGATGGACCTATACAAAATACACTTC CACACAAGCAGTGTCCTAAACCTAGGTAACATCTACTACCTGGATATGTTTTATGTGAAG
- .1120 GGCAAGCCAGAGTGGAAACCATAAGCACTGATGTGTGTAACAGAAAGGATGTGTATGATG CCGTTCGGTCTCACCTTTGGTATTCGTGACTACACACATTGTCTTTCCTACACATACTAC
- AGGGAGATTCTGGTGGACCTCTGGTTTATGATAATCATGACATCTGGTACATTGTAGGTA TCCCTCTAAGACCACCTGGAGACCAAATACTATTAGTACTGTAGACCATGTAACATCCAT
- TAGTAAGTTGGGGACAATCATGTGCACTTCCCAAAAAACCTGGAGTCTACACCAGAGTAA ATCATTCAACCCCTGTTAGTACACGTGAAGGGTTTTTTGGACCTCAGATGTGGTCTCATT
- CTAAGTATCGAGATTGGATTGCCTCAAAGACTGGTATGTAGTGTGGATTGTCCATGAGTT GATTCATAGCTCTAACCTAACGGAGTTTCTGACCATACATCACACCTAACAGGTACTCAA
- ATACACATGGCACAGAGCTGATACTCCTGCGTATTTTGTATTGTTTAAATTCATTTAC TATGTGTACCGTGTGTCTCGACTATGAGGACGCATAAAACATAACAAATTTAAGTAAATG
- TTTGGATTAGTGCTTTTGCTAGATGTCAAGAAGCCCTTCAGACCCAGACAAATCTAATAT AAACCTAATCACGAAAACGATCTACAGTTCTTCGGGAAGTCTGGGTCTGTTTAGATTATA
- CCTGAGGTGGCCTTTACATACGTAGGACCAAACCCTCTCTACCATGAGGGAAGAAGACAC GGACTCCACCGGAAATGTATGCATCCTGGTTTGGGAGAGATGGTACTCCCTTCTTCTGTG
- AGCAAATGACAGGACAGCTATTCCTTACTCACAAGGGAAACTGCTTGTGATACTTCCT TCGTTTACTGTCGTGGATAAGGAATGAGTGTTCCCTTTGACGAACACTATGAAGGA
- AATAAGATAAATAAGTGGTTTCCCTCAATTGAAGACAGGAACATCATTTTCCACAGGATA TTATTCTATTTATTCACCAAAGGGAGTTAACTTCTGTCCTTGTAGTAAAAGGTGTCCTAT
- TGAAGAGCTGCCAGTAATGCCAAAATCTTACCTCATATAATACCTGGAGCATGTGAGATT ACTTCTCGACGGTCATTACGGTTTTAGAATGGAGTATATTATGGACCTCGTACACTCTAA
- CTTCTAGTGAAAAAGAACAGTCTTCCCTGAAGACTCAGGGCTTCAACATTCTAGAACTGA GAAGATCACTTTTTCTTGTCAGAAGGGACTTCTGAGTCCCGAAGTTGTAAGATCTTGACT
- TAAGTGGACCTTCAGTGTGCAAGAATGGAGAAGCATGGGATTTGCATTATGACTTGAACT ATTCACCTGGAAGTCACACGTTCTTACCTCTTCGTACCCTAAACGTAATACTGAACTTGA
- GGGCTTATATCTAATAATACAGAGCACTATCACTAACCTCAACAGTTGACATTTTAAAAG CCCGAATATAGATTATGTCTCGTGATAGTGATTCGAGTTGTCAACTGTAAAATTTTC

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1930 1940 1950 1960 1970 1980 TTTTTAAATGTATCTGAACTTGCTGTTAACACAGTGTTATAACTCAAGCACTAGCTTCAG AAAAATTTACATAGACTTGAACGACAATTGTGTCACAATATTGAGTTCGTGATCGAAGTC 1990 2000 2010 2020 2030 2040 GAAGCATGTTGTTTAAGAGCTTTTTCTGATTATTCTTTAACAGCATCTTGCCATC CTTCGTACAACAACAATTCTTCGAAAAGACTAAATAAGAAATTGTCGTAGAACGGTAG 2070 2060 2080 2100

FIG. 1E